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Scoring table:

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1 (bases 1 to 436)

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2 Hillier L., Clark, N., Dubuque, T., Elliston, K., Hawkins, M., Holman, M., Kuchan, M., Kuchan, T., Le, M., Lennon, G., Marra, M., Holman, M., Kuchan, M., Kuchan, T., Le, M., Lennon, G., Marra, M., Parsons, J., Rifkin, L., Rohlfing T., Sacres, M., Tan, F., Trevaskis, E., Waterston, The Washing T., Sacres, M., Tan, F., Trevaskis, E., Waterston, The Washington University School of Medicine Washington University School of Medicine 4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108

Tel: 314 286 1800

Fax: 314 286 18
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B1279473 U1-K-DAO-
B1282803 U1 K-DAO-
B1279272 U1-K-DAO-
B1279513 U1-K-DAO-
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/dev_stage="19 weeks"
/lab_host="bH10B (ampicillin resistant)"
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/db_xref="GDB:1270181"
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normalization to a Cot " 5. Library constructed by M.Fatima Bonaldo. This library was constructed from the same fetus as the fetal lung library, Soares fetal lung
                                                                                                          Email: cst@watson.wustl.cdu
This clone is available royalty-free through LLNL: contart the This clone is available royalty-free through LLNL: contart the TMAGE Consortium (info@image.llnl.gov) for further intormation. Insert Length: 508 std Error: 0.00 seq primer: mob.REGA+ET High quality sequence stop: 394.
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                                     Washington University School of Medicine
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Hiller.L. Clark,N., Dubuque,T., Elliston,K., Hawkins,M., Holman,M., Mullaman,M., Kucaba,T., Le,M., Lennon,G., Marra,M., Parsons,J., Rikin,L., Rohlfinq,T., Soares,M., Tan,F., Trevaskis,E., Waterston,K., Williamson,A., Wohldmann,P., and Willson,R.
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Contact: Robert Strausberg, Ph.D.
Email: cwapbs-remail.nih.gov
Tissue Procurement: Chris Moskaluk, M.D., Ph.D., Michael R.
Fissue Procurement: Chris Moskaluk, M.D., Ph.D., Michael R.
Emmert-Buck, M.D., Ph.D. cDNA Library Preparation: Life
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Technologies, Inc. CoNA Library Arrayed by: Christa Pranqe, The
Genome Sequencing Center
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                    AW070916 420 bp mRNA linear EST 20-OCT-2000 xd31d01.xl NCI_CGAP_Br18 Homo sapiens cDNA clone IMAGE:2568385 3' similar to contains element TAR1 repetitive element ;, mRNA
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/clonc_lib."NCI_CGAP_Br18"
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primary tumors and two metastatic to ovary"
/lab_host."DH108"
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Mummalia; Eutheria; Primatus; Catarrhini; Hominidae; Homo.
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/db_xret-"taxon:9606"
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Dias Neto, E., García Correa, R., Verjovski-Almeida, S., Friones, M.K., Nagai, M.A., da Silva, W. Jr., Zaqo, M.A., Bordin, S., Costa, F.F., Goldman, G.H., Carvalho, A.F., Matsukuma, A., Baia, G.S., Simpson, D.H., Brunstein, A., deOliveira, P.S., Bucher, P., Jongeneel, C.V., O'Hare, M.J., Soares, F., Brentani, R.R., Reis, L.F., de Souza, S.J. and
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//dev_stage="Adult"
//dev_stage="Adult"
//note="Organ: breast; Vector: pucl8; Site_1: Smal; Site_2: Smal; A mini-library was made by cloning products derived from ORESTES PCR (U.S. Letters Patent application No. 196, 716 - Ludwig institute for Cancer Research) profiles into the PUC 18 vector. Reverse transcription of tissue mRNA and cDNA amplification were performed under low
                                                                                                                                                                                                            EST 10 JAN 2001
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This sequence was derived from the FAPESP/LICK Human Cancer Genome Project. This entry can be seen in the following URL (http://www.ludwig.org.br/scripts/gethtml2.pl?ti-IL26t2 11.2-BFO807-041000-176-G036t3-2000-10-046t4-1)
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1L2-BT0807-041000-176-G03 BT0807 Homo sapiens CDNA, mkNA sequence.
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Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo,
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Ludwig Institute for Cancer Research
Rua Prof. Antonio Prudente 109, 4 andar, 01509-010, Sao Paulo-SP,
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/db_xref="taxon:9606"
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NCI-CCAP http://www.ncbi.nlm.nih.gov/ncicgap.
NALIONAL Cancer Institute, Cancer Genome Anatomy Project (CGAP),
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This clone is available royalty-free through LLNL; contact the
IMAGE Consortium (into-timage.llnl.qov) tor further intormation.
Insert Length: 510—814 Error: 0.00
Seq primer: -40013 twd. fr from Amersham
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/sex "unknown"
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Contact: Robert Strausberg, Ph.D.
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Harrington, J.J., Sherf, B., Rundlett, S., Jackson, P.D., Perry, R., Cain, S., Leventhal, C., Thornton, M., Ramachandran, R., Whittinuton, J., Lerner, L., Costanzo, D., McElligott, K., Boozer, S., Mays, R., Smith, E., Veloso, N., Kilka, A., Hess, J., Cothren, K., Lo, K., Ottrubacher, J., Danzig, J. and Ducar, M. Cothren, K., Lo, K., Ottrubacher Creation of genome-wide protein expression libraries using random activation of gene expression
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//db_xrei-"Laxon:9606"
//clone_lib-"Athersys RACE Library"
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 464)
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RST21668 Athersys RAGE Library Homo sapiens cDNA, mkNA sequence.
BG202312
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185 actyctyrtycaaraactycgacractyctyctcctarractycaagraeryrtyrt (r)
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Fax: 216 361 9596
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Nat. Biotechnol. 19 (5), 440-445 (2001)
21227151
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/db_xref-"taxon:9606"
/clone_lib="Athersys RACE Library"
/cell_line="HT1080"
/note="See 'Creation of Genome-wide Protein Expression
/Ibrarles using Random Aclivation of Gene Expression
Nature Biotechnology, in press. Note that even though the
cell type indicated is HT1080, since a random activation
method was used, these sequence tags are not necessarily
                                                 urrariquaardarretetetetaccartqrigtaaayadattccaytttacccaaa 280
                                                                                                                                                                                                                                                               208
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 DB 10; Length 464;
                            Indels
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Athersys, Inc.
1201 Carnegie Ave, Cleveland, OH 44115, USA
TTC1: 216 431 9900
Fax: 216 361 9596
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В
Score 345.2; DB 10
Pred. No. 1.6e-78;
0; Mismatches 8,
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High quality sequence stop: 3
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73.38;
97.68;
           Hest Local Similarity 97.6
Matches 461; Conservative
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This sequence was derived from the FAPESP/LICR Human Cancer Genome
Project. This entry can be seen in the following URL
(http://www.ludwig.org.br/scripts/gethtml2.pl?tl=&t2=CM1-BT0738-250
Seq primer: puc 18 forward
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                                                                                                  Length 376;
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CM1-BT0738-250400-200-c12 BT0738 Homo sapiens CDNA,
BE092421
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Ludwig Institute for Cancer Research
Rua Prof. Antonio Prudente 109, 4 andar, 01509-010,
                                                                                              Score 331; DB 10; Length 3:
Pred. No. 7e-75;
0; Mismatches 15; Indels
       expressed in HT1080 under 103 c 67 g 104
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95.6%;
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Fax: +55-11-2707001
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Location/Qualitiers
                             stop:
    Fax: 216 361 9596
Email: scain@athersys.com
High quality sequence stop
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                                             /Organism-"Homo sapiens"

/db_xref "Laxon: 4606"
/clone_lib-"Hru718#

/dev_stage."Adult"
/note-"Organ: breast; Vcctor: pucl8; Site_1: Sma1; Site_2:
Sma1; A min: library was made by cloning products derived
from ORESTES PCR (U.S. Letters Patent application No. 196
.716 - Ludwig institute for Cancer Research) profiles
into the pUC 18 vector. Reverse transcription of tissue
mink and cDNA amplification were performed under low
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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AUTHORS
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                                                                                                                                   Libraries using Random Activation of Gene Expression., Nature Biotechnology, in press. Note that even though the cell type indicated is HT1080, since a random activation method was used, these sequence tags are not necessarily expressed in HT1080 under normal circumstances.*
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I (bases 1 to 403)
Hillier,L., Clark,N., Dubuque,T., Elliston,K., Hawkins,M., Holman,M., Hultman,M., Kucaba,T., Le,M., Lennon,G., Marra,M., Parsons,J., Rifkin,L., Rohlfing,T., Soares,M., Tan,F., Trevaskis,E., Waterston,The Washu Werck EST Project
Unpublished (1995)
/organism="Momo sapiens"
/db_xref="taxon:9606"
/cblone_lib="Athersys RACE Library"
/cell_line="HT1080"
/note="See 'Creation of Genome-wide Protein Expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    304
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Washington University School of Medicine
4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108
Tel: 314 286 1800
Fax: 314 286 1810
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97.1%; Pred. No. 2.1e-69;
ive 0; Mismatches 8; Indels
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Best Local Similarity
Matches 313; Conserv
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//lab_host="NHIOB" (ampicillin resistant)"
//lab_host="NHIOB" (ampicillin resistant)
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           Final in carractions of the property of the primer. Might property of the primer of the primer. Might property of the primer of the primer. Might property of the primer of the primer. Might property of the primer of the primer. Might property of the primer of the primer. Might property of the primer of the primer of the primer of the primer of the primer. Might property of the primer of 
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6
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/db_xref="CbB:572980"
/db_xref="taxon:9906"
/clone="IMAGE:158921"
/rlone_11b-"Soares breast 2NbHBst"
/scx="Female"
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Pred. No. 3.7e-69;
0; Mismatches 15;
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Email: estewatson.wustl.edu
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Aurora Biosciences Corp.

Aurora Biosciences Corp.

1000 Torreyana Road, San Diego, CA 92121, USA
Tel: 8584048736

Fax: 8584046719

Email: henkelg@aurorabio.com
Pools of cells were isolated from a GenomeScreen(TM) library. The
library of cells was generated by retroviral integration of a gene
tagging element consisting of: 1) A promoterless beta-lactamase
proceeded by a splice acceptor as a reporter for gene expression;
2) A promoter driving neomycin resistance followed by a splice
donor to trap downstream exons. 3' RACE from neomycin gene was
performed using total RNA from isolated pools. Output was shotyun
cloned in pAmp-1 and used to transform DH5-alpha competent
bacteria. 5' ends of reported sequences were immediately proceded
Class: exon-trapped.
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostom); Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. (bases 1 to 314)
Henkel,G., Liyanage,M., Pratt,E., Huang,D., Riley,M., Bernardino,A., Durick,K. and Pollok,B.

Exon trap tags from a HL-60 GenomeScreen(TM) Library
Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /note="Organ: peripheral blood; Vector: pAmp-1; i' RACE or tcotal RNA from genetrap pools; shotgun clone in pAmp-1 and used to transform DHS-alpha competent bacteria."
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /tissue_type="acute promyelocytic leukemia"
/cell_type="promyeloblast"
/cell_line="HL-60"
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/db_xref="taxon:9606"
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                                                                                                                                                                                      Contact: Greg Henkel
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/db_xref-"taxon:9606"
/db_xref-"taxon:9606"
/clone_lib_rathersys RACE Library"
/cell_line."HT1080*
/note-"see 'Creation of Genome-wide Protein Expression
Libraries using Random Activation of Gene Expression
Libraries using Random Activation of Gene Expression
(Libraries using Random Activation of Gene Expression
Libraries using Random Activation of Gene Expression
(Libraries using Random Activation of Gene Expression)
(Nature Biotechnology, in press. Note that even though the
Cell type Indicated is HT1080, since a random activation
method was used, these sequence tags are not necessarily
expressed in HT1080 under normal circumstances."
                                    EST 21-APR-2001
                                                                                                                                                                                      Harrington, J.J., Shert, B., Rundlett, S., Jackson, P.D., Perry, R., Cain, S., Leventhal, C., Thornton, M., Ramachandran, R., Whittington, J., Lerner, L., Costanzo, D., McElligott, K., Boozer, S., Mays, R., Smith, E., Veloso, N., Klika, A., Hess, J., Cothren, K., Lo, K., Offenbacher, J., Danziq, J. and Ducur, M. Credion of genew-wide protein expression libraries using random activation of gene expression.
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                                                                                                                            Homo sapiens
Eukaryota; Metazoa: Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                 HG202313 143 bp mkNA linear EST 21-APR-2
KST21669 Athersys KAGE Library Homo Sapiens CDNA, mkNA Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                    High quality sequence stop: 343.
                                                                                                                                                                                                                                                                                                                                                     Athersys, Inc.
3201 Carneque Ave, Cleveland,
TPI: 216 431 9900
Fax: 216 361 9596
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Location/Qualitiers
                                                                                                                                                                                                                                                                                                                                                                                                                     scain@athersys.com
                                                                             BG202313.1 G1:13724000
                                                                                                                                                                                                                                                                                                                                        Contact: Scott J. Cain
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93.3%;
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Email: asimpsoneludwig.org.br
This sequence was derived from the FAPESP/LICR Human Canter Genome
Project. This entry can be seen in the following URL
(http://www.ludwig.org.br/scripts/gethtml2.pl?tl-QV0&L2-QV0-BT0103-
Seq primer: puc 18 forward
High quality sequence start: 7
High quality sequence stop: 313.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone_lib="BT00103"
/dev_stage="Adult"
/dev_stage="Adult"
/note="Organ: breast; Vector: puc18; Site_1: Sma1; Site_2: Sma1; A mini-library was made by cloning products derived from ONESTES PCR (U.S. Letters Patent application No. 196. 716 - Ludwig Institute for Cancer Research) profiles into the pUC 18 vector. Reverse transcription of tissue mank and clonk amplification were performed under low
        EST 16-N:IV-1999
AW176044 313 bp mRNA linear ESF 16-New-199
OVO-BF0103-270899-008-112 BT0103 Homo sapiens CDNA, mKNA sequence.
                                                                                                                                               Eukaryota, Metazoa, Chordata, Craniala, Vertebrata, Euloleostomi,
Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
                                                                                                                                                                                                                                                                                               Laboratory of Cancer Genetics
Ludwig Institute for Cancer Research
Rua Prof. Antonio Prudente 109, 4 andar, 01509-010, Sao Panlo-SP,
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Pred. No. 2.3e-65;
0; Mismatches 5; Indels
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The FAPESP/LICR Human Cancer Genome Project
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                                                                                                                                                                                                                                                                                  Contact: Simpson A.J.G.
                                                             AW176044.1 G1:6442U81
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98.1%;
                                                                                                                                                                                         (bases 1 to 313)
                                                                                                                                                                                                                                                                                                                                                                                   Tel: +55-11-2704922
                                                                                                                                                                                                                                                                                                                                                                                                       Fax: +55-11-2707001
                                                                                                                                                                                                                                                         Unpublished (1999)
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us-09-525-361a-23.rst

DEFINITION

ACCESS 10N

VERSION KEYWORDS

ORGANISM

SOURCE

REFERENCE

AUTHORS

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RESULT

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AA340069

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Fax: +55-11-2707001
Email: asimpson@ludwig.org.br
This sequence was derived from the FAPESP/LICR Human Concer Genome
Project his entry can be seen in the following URI.
(http://www.ludwig.org.br/scripts/gethtml2.pl?tl~6t2.cMl·HT07.88-250
400-199-bl16t3=2000-04-25&t4=1)
Seq primer: puc 18 forward
High quality sequence start: 4
High quality sequence start: 4
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93 c 1 g 82 t 1 others
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Laboratory of Cancer Genetics
Ludwig Institute for Cancer Research
Rua Prof. Antonio Prudente 109, 4 andar, 01509-010, Sao Paulo-SP,
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Proc. Natl. Acad. Sci. U.S.A. 97 (7), 3491-3496 (2000)
20202663
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Pred. No. 5.3e-62;
0; Mismatches 5; Indels
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/organism≈"Homo sapiens"
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/clone_lib="HT0738"
/dev_stage="Adult"
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Best Local Similarity 97.8%;
Matches 306; Conservative
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/tissue_type="epithelium"
/tissue_type="epithelium"
/dev_stage="fetus"
/note="Oran: skin; Vector: pBluescript SK-; Site_l: EcoRI: Site_2: XhoI"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Initial assessment of human gene diversity and expression patterns based upon 83 million nucleotides of cDNA sequence Nature 377 (6547 Suppl), 3-174 (1995)
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For clone availability, additional sequence and expression
information related to this EST, please click the TIGR Human Gene
Index (http://www.tigr.org/tdb/hqi/hgi.html)
Seq primer: M13 Reverse.
                                                                                                                                                                                                                                                                                                                                                                                                               Euteleostomi;
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Enkaryota; Metazoa; Chordata; Craniata; Vertebrata; Eutele
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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9712 Medical Center Drive, Kockville, MD 20850 USA
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/db_xref-"taxon:9606"
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AA340069.1 GI:1992307
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Contact: Kerlavage, AR
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Search completed: October 5, 2002, 05:07:20 Job time: \$805 sec 21, Appl 21, Appl 21, Appl 5273901

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Pred. No. 8.6e-126;
0; Mismatches 6;
                          US-08-770-379-20
US-08-770-379-20
US-08-77-669A-20
US-08-718-661-1
US-08-82-170A-20
US-09-273-565-20
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APPLICANT: Dumas Milne Edwards, Jean-Baptiste
APPLICANT: Dumas Milne Edwards, Jean-Baptiste
APPLICANT: Duclert, Aymeric
APPLICANT: Bougueleret, Lydie
TITLE OF INVENTION: Complementary DNAS
FILE REFERENCE: GENSET.021A
CURRENT APPLICATION NUMBER: US/09/247,125
EARLIER APPLICATION NUMBER: 60/074,121
EARLIER APPLICATION NUMBER: 60/081,563
EARLIER APPLICATION NUMBER: 60/080;
EARLIER PELING DATE: 1998-04-13
EARLIER FILING DATE: 1998-06-10
EARLIER FILING DATE: 1998-06-10
EARLIER FILING DATE: 1998-08-10
EARLIER FILING DATE: 1998-10-04
NUMBER OF SEQ ID NOS: 182
SOPTWARE: PATENT.PM
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INFORMATION: seq LVLLGVSIFLVSA/QN
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                                                                                                                                                                                                                                                                                                                                             Sequence 41, Application US/09247155A Patent No. 6312922 GENERAL INFORMATION:
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98.58;
  Best Local Similarity 98.5
Matches 463; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: DNA
ORGANISM: Homo sapiens
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NAME/KEY: polyA_signal
LOCATION: 462..467
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LOCATION: 68..124
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    μαμαμοσφοσφοσφορώ
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Sequence 1, Ap
Sequence 3, Ap
Sequence 3, Ap
Sequence 29, P
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Sequence 4, A
Sequence 5, A
Sequence 5, A
Sequence 2, A
Sequence 6, A
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Sequence 7,
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2: /cgnl__6/ptodata/1/ina/5B_COMB.seq:*
3: /cgnl__6/ptodata/1/ina/6A_COMB.seq:*
4: /cgnl__6/ptodata/1/ina/FG_TUS_COMB.seq:*
5: /cgnl__6/ptodata/1/ina/FG_TUS_COMB.seq:*
6: /cgnl__6/ptodata/1/ina/PCTUS_COMB.seq:*
     GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd.
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US-08-20-9802B-3

US-08-20-859-29

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US-08-20-802B-4

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US-09-359-081-2

US-09-10-647-1

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                                                                                                                                                                                                                                                                 hits satisfying chosen parameters:
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                                                                                     October 5, 2002, 03:59:20;
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Listing first 45 summaries
                                                        OM nucleic - nucleic search, using sw model
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Gapop 10.0 , Gapext 1.0
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TITLE OF INVENTION: Diagnostic Method and Kit for Neuropsychiatric Diseases
TITLE OF INVENTION: Using Trinucleotide Repeats Sequence
FILE REPERENCE: 1942/36
CURRENT FILING DATE: 1999-02-22
EARLIER APPLICATION NUMBER: KR 98-6,278
EARLIER FILING DATE: 1996-02-26
NUMBER OF SEQ ID NOS: 3
SOFTWARE: Wordberfect 6.1/Windows
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                                                                                                                                                                                                                                                                                                                                                                                                Length 397;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-09-135-994-1/c
: Sequence 1, Application US/09135994A
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Patent No. 6124100
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Best Local Similarity
Matches 110; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
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US-09-253-691-3
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159 atgatgaagcccctgatgctgaaaccactgctgctgcaacaactgcgaccactgctgctc 218
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APPLICANT: Ranum, Laura P.W.
APPLICANT: Chung, Ming-yi
APPLICANT: Coghbi, Huda Y.
TITLE OF INVENTION: Gene Sequence for Spinocerebellar Ataxia
Patent No. 5741645
TITLE OF INVENTION: Type 1 and Method for Diagnosis
NUMBER OF SEQUENCES: 47
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                         Length 477;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SSEE: Mueting, Raasch, Gebhardt & Schwappach,
P: 119 No. 5741645th Fourth Street, Suite 203
Minneapolis
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Ranum et al.
TITLE OF INVENTION: SCA7 GENE AND METHODS OF USE
FILE REFERENCE: University of Minnesota
CURRENT APPLICATION NUMBER: US/09/135, 994A
CURRENT FILING DATE: 1998-08-18
EARLIER FILING DATE: 1997-08-19
NUMBER OF SEQ ID NOS: 14
SOFTWARE: PATENTIN Ver: 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA: PAPLICATION NUMBER: US/08/469,802B FILING DATE: 06-JUN-1995 CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                  13.9%; Score 65.4; DB 4; 63.9%; Pred. No. 1.9e-10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        180 CTGCTGCTGCTGCTGCTGCTGCTGCTGCTGT 146
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; Sequence 3, Application US/08469802B
; Patent No. 5741645
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: MUELING, AND M.
REGISTRATION NUMBER: 33,977
REFERENCE/DOCKET NUMBER: 11
TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION:
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Best Local Similarity 63.97
Matches 99; Conservative
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                                                                                                                                                                                                                                                                                                           ; ORGANISM: Homo sapiens
US-09-135-994-1
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APPLICANT: Ranum, Laura P.W.
APPLICANT: Chung, Ming-yi
APPLICANT: Zoghbi, Huda Y.
TITLE OF INVENTION: Gene Sequence for Spinocerebellar Ataxia
Patent No. 5834183
                                                                                                                                         13.8%; Score 65; DB 1; Length 234; 58.5%; Pred. No. 1.7e-10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEE: Mueting, Raasch, Gebhardt & Schwappach, P.A.: P.O. Box 581415
Minneapolis
                                                                                                                                                                               80; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: Type 1 and Method for Diagnosis NUMBER OF SEQUENCES: 85
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM PC Competible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/267,803B
FILING DATE: 28-JUN-1994
CLASSIFICATION: 435
                                                                                                                                                                            0; Mismatches
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Patent No. 5834183
GENERAL INFORMATION:
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NAME: MCCOFFMACK, Myra H.
RECISTRATION NUMBER: 36.002
REFERENCE/DOCKET NUMBER: 110
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELECOMMUNICATION INFORMATION:
TELEPHONE: 612-305-1217
TELEFAX: 612-305-1228
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INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
LENGTH: 234 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLLGY: linear
MOLECULE TYPE: DNA
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                                                                                                                                                       Best Local Similarity 58.5
Matches 113; Conservative
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TYPE: nucleic acid
STRANDEDNESS: single
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ADDRESSEE: Mueting,
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MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                        241 ttctaccactgct 253
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MOLECULE TYPE: DNA
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US-08-267-803B-3/c
                                                                                    US-08-469-802B-3
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                                                                                                                                           Query Match
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Patent No. 563135
GENERAL INFORMATION:
GENERAL INFORMATION:
TITLE OF INVENTION: Diagnostics, Therapeutics and Methods for TITLE OF INVENTION: Detection and Treatment of Acute Leukemias
TITLE OF INVENTION: Resulting from Chromosome Abnormalities in the NUMBER OF SEQUENCES: 44
                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
ADDRESSEE: Woodcock Washburn Kurtz Mackiewicz & No. 5633135ris
STREET: One Liberty Place - 46th Floor
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       Length 234;
                                         80; Indels
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   Score 65; DB 2; I
Pred. No. 1.7e-10;
0; Mismatches 80;
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FILING DATE: 14 MAY 1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/07/971,094
FILING DATE: 30-0CT-92
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/07/888,830
FILING DATE: 27-MAY-92
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/07/805,093
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/320,559
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ATTORNEY/AGENT IN-DEC-91
ATTORNEY/AGENT INFORMATION:
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 13.8%;
58.5%;
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TELEFACK: (215) 568-3439
INFORMATION FOR SEQ ID NO: 29:
SEQUENCE CHARACTERISTICS:
LENGTH: 3376 base pairs
Query Match 13.8
Best Local Similarity 58.5
Matches 113; Conservative
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PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ZIP: 19103
COMPUTER READABLE FORM:
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OPERATING SYSTEM: E
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US-08-320-559-29/c
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Sequence 29, Application PC/TUS9404496
GENERAL INFORMATION:
APPLICANT: Croce, Carlo
APPLICANT: Canaani, Eli
TITLE OF INVENTION: Diagnostics, Therapeutics and Methods
TITLE OF INVENTION: For Detection and Treatment of Acute Leukemias
TITLE OF INVENTION: Resulting from Chromosome Abnormalities in the All-1
NUMBER OF SEQUENCES: 86
CORRESPONDENCE ADDRESS:
ADDRESSEE: Woodcock, Washburn, Kurtz, Macklewitz 6
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Best Local Similarity 57.4%; Pred. No. 8.5e-10;
Matches 117; Conservative 0; Mismatches 87;
                                                                                                                                                         APPLICATION NUMBER: US 07/888,839
FILING DATE: 27-MAY-1992
PRIOR APPLICATION DATE: US 07/805,093
FILING DATE: 11-DEC-1991
ATTORNEY/AGENT INFORMATION:
NAME: Deluca Esq., Mark
REGISTRATION NUMBER: 33,229
REFERENCE/DOCKET NUMBER: TJU-1262
TELECOMMUNICATION INFORMATION:
TELEPHONE: (215,568-3100
TELE
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                                                                           APPLICATION NUMBER: US 07/971,094
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             223 cactgcaaccaccgctgcttctac 246
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                                                                                                    FILING DATE: 30-OCT-1992
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 3376 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
                                               PRIOR APPLICATION DATA:
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy of
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FOCATION: 196..1902
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Patent No. 6040140
GENERAL INFORMATION:
Canaani, Eli
TTLE OF INVENTION: Diagnostics, Therapeutics and Methods
TITLE OF INVENTION: Diagnostics, Therapeutics and Methods
TITLE OF INVENTION: Resulting from Chromosome Abnormalities in the All-1 Region
NUMBER OF SEQUENCES: 94
CORRESPONDENCE ADDRESS:
ADDRESSEE: Woodcock, Washburn, Kurtz, Mackiewitz &
ADDRESSEE: Woodcock, Washburn, Kurtz, Mackiewitz &
CORREST: One Liberty Place, 46th floor
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                                                                                                                                                                                                                                                                                                   13.8%; Score 64.8; DB 1; Length 3376; llarity 57.4%; Pred. No. 8.5e-10; Conservative 0; Mismatches 87; Indels 0
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPALIALE
COMPUTER: IBM PC COMPALIALE
COMPUTER: PATENTION DATA:
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/545,860D
FILING DATE: 07-MAR-1996
CLASSIFICATION: 435
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APPLICATION NUMBER: PCT/US94/04496
FILING DATE: 22-APR-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/US92/10930
FILING DATE: 09-DEC-1992
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US 08/327,392
FILING DATE: 19-OCT-1994
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APPLICATION NUMBER: US 08/320,559
FILING DATE: 11-0CT-1994
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                                                                                       MOLECULE TYPE: DNA (genomic) FEATURE:
TYPE: nucleic acid
STRANDEDNESS: double
TOPOLOGY: linear
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APPLICATION NUMBER:
                                                                                                                                                                                  196..1902
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USA
                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
Matches 117; Conserv
                                                                                                                                                      CDS
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                                                                                                                                                   ; NAME/KEY:
; LOCATION:
US-08-320-559-29
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                                             101 gcccagaatccgacaacagctgctccagctgacacgtatccagctactggtcctgctgat 160
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                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
APPLICANT: Orr, Harry T.
APPLICANT: Ranum, Laura P.W.
APPLICANT: Chung, Ming-yi
APPLICANT: Coghbi, Huda Y.
TITLE OF INVENTION: Gene Sequence for Spinocerebellar Ataxia
TITLE OF INVENTION: Type 1 and Method for Diagnosis
NUMBER OF SEQUENCES: 47
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  13.5%; Score 63.4; DB 1; Length 168; ilarity 63.4%; Pred. No. 4.4e-10; Conservative 0; Mismatches 56; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADDRESSEE: Mueting, Raasch, Gebhardt & Schwappach, STREET: 119 No. 5741645th Fourth Street, Suite 203 CITY: Minneapolis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/469,802B
FILING DATE: 06-JUN-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               REFERENCE/DOCKET NUMBER: 110.00030101
ELECOMMUTCATION INFORMATION:
TELEPHONE: 612-305-1217
TELEFAX: 612-305-1225
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                       Sequence 4, Application US/08469802B
Patent No. 5741645
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 33,977
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ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH: 168 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Mueting, Ann M. REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
Matches 97; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; MOLECULE TYPE:
US-08-469-802B-4
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                                                                                                                                                                                                                                         269 gtt 271
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APPLICANT: TSUJI, Shoji
APPLICANT: TSUJI, Shoji
APPLICANT: SANDEI, Kazujiro
APPLICANT: SANDEI, Kazujiro
TITLE OF INVENTION: Method for Diagnosing Spinocerebellar Alaxia Type 2 and
TITLE OF INVENTION: Primers Therefor
TITLE OF INVENTION: Primers Therefor
TITLE OF INVENTION: Primers Therefor
CURRENT APPLICATION NUMBER: US/09/043,303
CURRENT FILING DATE: 1998-05-18
EARLIER FILING DATE: 1996-07-18
NUMBER OF SEQ ID NOS: 17
SOFTWARE: Patentin Ver. 2.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 844 CACTITTAIGITCTCTGGAGTCTTIAGAAGGTTTTTCCTTGTGCTCCTTCATTAATTTGT 785
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 3376;
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Pred. No. 2.1e-10;
): Mismatches 74; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      87; Indels
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SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            13.8%; Score 64.8; DB 5;
57.4%; Pred. No. 8.5e-10;
tive 0; Mismatches 87;
                                                                                   ATTORNEY/AGENT INFORMATION:
NAME: DeLuca ESq., Mark
REGISTRATION NUMBER: 33,229
REFERENCE/DOCKET NUMBER: TJU-1242
TELECOMMUNICATION INFORMATION:
TELEPHONE: (215) 568-3100
TELERAX: (215) 568-3100
TELERAX: (215) 568-3439
INFORMATION FOR SEQ ID NO: 29:
SEQUENCE CHARACTER ISTICS:
                                   APPLICATION NUMBER: PCT/US94/04496
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             223 cactqcaaccaccgctgcttctac 246
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: Sequence 7, Application US/09043303
: Patent No. 6251589
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MOLECULE TYPE: DNA (genomic)
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59.6%;
                                                                                                                                                                                                                                                                LENGTH: 3376 base pairs
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Best Local Similarity 59.6
Matches 109; Conservative
                                                                                                                                                                                                                                                                              TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                               ; NAME/KEY: CDS
; LOCATION: 196..1902
PCF-US94-04496-29
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                                                                           CLASSIFICATION:
                                                     FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQ ID NO 7
LENGTH: 203
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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Patent No. 5834183
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE: 06-JUN-1995
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: MUELING, Ann M.
REGISTRATION NUMBER: 33,977
REJENENCE/OCKET NUMBER: 110
TELECOMMUNICATION INFORMATION:
TELEPHONE: 612-305-1217
TELEFAX: 612-305-1225
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TYPE: nucleic acid
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MEDIUM TYPE: Floppy
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                               Patent No. 5741645
GENERAL INFORMATION:
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US-08-267-803B-5/C
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161 gatgaagccctgatgctgaaccactgctgctgcaacaactgcgaccactgctgctcct 220
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                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Orr, Harry T.
APPLICANT: Ranum, Laura P.W.
APPLICANT: Chung, Minq-yi
APPLICANT: Zoghbi, Huda Yi
TITLE OF INVENTION: Gene Sequence for Spinocerebellar Ataxia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 168;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      E: Mueting, Raasch, Gebhardt & Schwappach, P.A.
P.O. Box 581415
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Patent No. 5834183
TITLE OF INVENTION: Type 1 and Method for Diagnosis NUMBER OF SEQUENCES: 85
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/267,803B
FILING DATE: 28-JUN-1994
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         13.5%; Score 63.4; DB 2; 63.4%; Pred. No. 4.4e-10; ative 0; Mismatches 56;
                                                                      221 accactgcaaccaccyctgcttctaccactgct 253
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               43 GCTGCTGCTGCTGCTGCTGCTGCTGCT 11
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 612-305-1217
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                  Sequence 4, Application US/08267803B
FALENT NO. 5843183
GENERAL INFORMATION:
APPLICANT: Orr, Harry T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-08-469-8028-5/c Sequence 5, Application US/08469802B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ZIP: S5458-1415
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEFAX: 612-305-1220
[NFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
LENGTH: 168 base pairs
TYPE: nucleic acid
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NAME: MCCOrmack, Myra H.
REGISTRATION NUMBER: 36,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity 63.49
Matches 97; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
ADDRESSEE: Mueting, I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CITY: Minneapolis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MOLECULE TYPE: DNA
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101 gcccagaatccgacaacagctgctccagctgacacgtatccagctactggtcctgctgat 160
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APPLICANT: Ranum, Laura P.W.
APPLICANT: Chung, Ming-yi
APPLICANT: Zoghibi, Huda Yi
APPLICANT: Zoghibi, Huda Yi
TITLE OF INVENTION: Gene Sequence for Spinocerebellar Ataxia
Patent No. 5741645
TITLE OF INVENTION: Type 1 and Method for Diagnosis
NUMBER OF SEQUENCES: 47
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
APPLICANT: Orr, Harry T.
APPLICANT: Ranum, Laura P.W.
APPLICANT: Chung, Ming-Yi.
APPLICANT: Zoghbi, Huda Y.
TITLE OF INVENTION: Gene Sequence for Spinocerebellar Ataxia
TITLE OF INVENTION: Type 1 and Method for Diagnosis
NUMBER OF SEQUENCES: 85
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             13.5%; Score 63.4; DB 1; Length 171; 63.4%; Pred. No. 4.4e-10; Live 0; Mismatches 56; Indels
                                                                                                                                                                                                                               ADDRESSEE: Mucting, Raasch, Gebhardt & Schwappach, STREET: 119 No. 5741645th Fourth Street, Suite 203
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/469,802B
FILING DATE: 06-JUN-1995
CLASSIFICATION: 435
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ATTORNEY/AGENT INFORMATION:
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TYPE: nucleic acid
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Best Local Similarity 63.4
Matches 97; Conservative
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CITY: Minneapolis
STATE: MN
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                                                                                                      CLASSIFICATION:
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                                                                                                                                                                                                                                                                                                           TYPE: nucleic STRANDEDNESS:
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US-08-267-803B-2/C
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Patent No. 5741645
GENERAL INFORMATION:
APPLICANT: Orr. Harry T.
APPLICANT: Chung, Ming-yi
APPLICANT: Zoghbi, Huda Y.
TITLE OF INVENTION: Gene Sequence for Spinocerebellar Ataxia
Patent No. 5741645
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 63.4; DB 2; Length 171;
Pred. No. 4.4e-10;
0; Mismatches 56; Indels (
CORRESPONDENCE ADDRESS:
ADDRESSE: Mueting, Raasch, Gebhardt & Schwappach, P.A.
STREET: P.O. Box 581415
CITY: Minneapolis
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                                                                                                COMPUTER READABLE FORM:
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: 1BM PC compatible
UPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PETCHILL Release #1.0, Version #1.25
CUSRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/267,803B
FILING DATE: 28-JUN-1994
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: Type 1 and Method for Diagnosis NUMBER OF SEQUENCES: 47 CORRESPONDENCE ADDRESS:
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NAME: MCCOTMACK, Myra H.
REGISTRATION NUMBER: 36,602
REFERENCE/POCKET NUMBER: 110.0
TELECOMMUNICATION INFORMATION:
TELEFAX: 612-305-1217
TELEFAX: 612-305-1217
TELEFAX: 612-305-1217
SEQUENCE CHARACTERISTICS:
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 13.58;
Best Local Similarity 63.48;
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TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               97; Conservative
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US-08-267-8038-5
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                                                                       STATE: N
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101 gcccagaatccgacaacagctgctccagctgacacgtatccagctactggtcctgctyat 160
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APPLICANT: Ranum, Laura P.W.
APPLICANT: Chung, Ming-yi
APPLICANT: Zoghbi, Huda Y
TITLE OF INVENTION: Gene Sequence for Spinocerebellar Ataxia
Patent No. 5834183
TITLE OF SEQUENCES: 85
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   13.5%; Score 63.4; DB 1; Length 195; 63.4%; Pred. No. 4.8e-10; tive 0; Mismatches 56; Indels
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STREET: P.O. Box 581415
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
FILLING DATE: 05-JUN-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/267,803B
FILING DATE: S8-JUN-1994
CLASSIFICATION: 435
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      70 GCTGCTGCTGCTGCTGCTGCTGCTGCT 38
                                                                                                                                                          NAME: MUETING, AND M.
REGISTRATION NUMBER: 33,977
REFERENCE/DOCKET NUMBER: 110.00030101
TELECOMMUNICATION INFORMATION:
TELEPHONE: 612-30-127
TELEPHONE: 612-30-125
INFORMATION FOR SEQ 1D NO: 2:
SEQUENCE CHARACTERISTICS:
                                    US/08/469,802B
06-JUN-1995
NI: 435
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NAME: MCCOrmack, Myra H.
REGISTRATION NUMBER: 36,602
REFERENCE/DOCKET NUMBER: 110.00030120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ZIP: 55458-1415
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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; Patent No. 5834183
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| TELECOMMUNICATION INFORMATION:
| TELEPAX: 612-305-1217 |
| TELEPAX: 612-305-1218 |
| TELEPAX: 612-305-1228 |
| TELEPAX: 612-305-1228 |
| TELEPAX: 612-305-1228 |
| TELEPAX: 612-305-1228 |
| SEQUENCE CHRACTERISTICS: |
| SEQUENCE CHRACTERISTICS: |
| LENGTH: 195 base pairs |
| TYPE: nucleic acid |
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Search completed: October 5, 2002, 05:08:26 Job time: 4146 sec

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15-MAR-1999;
12-NOV-1999;
12-NOV-1999;
15-NOV-1999;
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3815.365 Million cell updates/sec
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DNA encoding human
cDNA sequence of h
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SIDSI/gcgdata/geneseq/geneseqn-embl/NA1980 DAT:*
SIDSI/gcgdata/geneseq/geneseqn-embl/NA1981.DAT:*
SIDSI/gcgdata/geneseq/geneseqn-embl/NA1981.DAT:*
SIDSI/gcgdata/geneseq/geneseqn-embl/NA1981.DAT:*
SIDSI/gcgdata/geneseq/geneseqn-embl/NA1985.DAT:*
SIDSI/gcgdata/geneseq/geneseqn-embl/NA1985.DAT:*
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SIDSI/gcgdata/geneseq/geneseqn-embl/NA1999.DAT:*
SIDSI/gcgdata/geneseq/geneseqn-embl/NA1991.DAT:*
                                                                                                                      October 5, 2002, 04:39:04; Search time 211.95 Seconds
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GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd.
                                                                                                                                                                                                                                                                                                                                                                                          Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                       1736436 seqs, 858457221 residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SUMMARIES
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Maximum Match 100%
Listing first 45 summaries
                                                                               OM nucleir - nucleic search, using sw model
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AAF85500
AAX84202
AAC79431
AAX00704
AAX400644
AAC79470
AAI67224
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Gapop 10.0 , Gapext 1.0
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Maximum DB seq length: 2000000000
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Match Length DB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Title:
Perfect score:
                                                                                                                                                                                                                                                                                           Scoring table:
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                                                                                                                                                                                                                                                                                                                                                       Searched:
                                                                                                                                                                                                                                                  Sequence:
                                                                                                                         Run on:
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	16	446.2			473		AAD18674		Human BS106	o genera
	17	443			471		AAZ65099		Ø	ound pro
	18	443			471		AAF44245		Human PRO1	160 (UNO
	19	443			471		AAC97492			
	075	2.2.2			468		AAF44870		Human breast	st cance
	22	900			300		AAV 51990		BS106 poly	polynucleoti
	23	3008			308		AAD18670		HUMBO CION	093900
	24	298.8			501		AAS87721		DNA encoding povel	d povel
	25	286.8		6.	292		AAD18671		Human BS106	g dene-s
	56	271.4		9.	273		AAA54121		Breast cancer	١,
	27	255.6		.3	264		AAF93399			Ing SRT
	28	255.6		۳.	266		AAF93398		cDNA encoding	ing SRT
	53	247.2			291		AAF93397		cDNA encod	ing SRT
	30	202.8		~ .	229		AAV31989		BS106 polyr	ucleoti
	7 C	207.8			229		AAX55578		BS106 clone	6 clone 166288
	7 .	180.4		0.	707		AAD18669		Human BS10	gene-s
	n c	101.4		٠ ا	767		AAV31991		BS106 polyr	ucleoti
	+ t	181.4			197		AAX3558U		BS10b clone	clone 120981
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U	37	70.2			397		AAX89891		Spinocorpholist at	ig novel
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AAF85500 standard; cDNA; 471
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                                                                                                                              Screening drug candidates for their ability to modulate breast cancer by contacting the drug to a cell expressing an expression profile gene and determining modulation of expression of the gene
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  tgctccagctgacacgtatccagctactggtcctgctgatgatgaagcccctgatgctga 180
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                Comprise adding a drug candidate to a cell that expresses a protein selected from BCH1, BCA2, BCN7, BCN1, BCN5, BCO2, BCO2, BCR2, BCN2 and BCY3 or their fragments and determining the effect of the drug on the expression of those proteins. Antibodies to breast cancer speeds (specifically BCH1 or its fragment (BCH1p1 or BCH1p2)) are useful for inhibiting and treating breast cancer in individuals who are non-responsive to anti-ocstrogen and positive for oestrogen receptor. Compositions comprising BCH1 or a nucleic acid encoding BCH1 are useful for eliciting an immune response in an individual. The antibodies are also useful for the diagnosis and prognosis of breast cancer and for screening compositions which modulate the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               aaccactgctgctacaacaactgcgaccactgctcctaccactgcaaccaccgctgc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ttotaccactgotcgtaaagacattccagttttacccaaatgggttggggatctcccgaa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     tggtagagtgtgtccctgagatggaatcagcttgagtcttctgcaattgggtcacaacta
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                                                                                                                                                                                                                                                                                                                                                                            DB 21; Length 471;
                                                                                                                                                                                          New methods for screening drug candidates are described which
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   breast cancer phenotype. The method allows rapid and simple detection of lymph node metastases.
                                                                                                                                                                                                                                                                                                                                                                                                Indels
                                                                                                                                                                                                                                                                                                                                             Sequence 471 BP; 108 A; 133 C; 88 G; 142 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                     3.4e-119;
                                                                                                                                                                                                                                                                                                                                                                                    Pred. No. 3.4e
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                           Score 469.4;
                                                                                                                                                                       Disclosure; Fig 32; 258pp; English.
99US-0440676.
99US-0440677.
99US-0450810.
99US-0453137.
                                                            (EOSB-) EOS BIOTECHNOLOGY INC
                                                                                                                                                                                                                                                                                                                                                                         99.78;
                                         08-MAR-2000; 2000US-0453137.
                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity 99.8 Matches 470; Conservative
                                                                                                  WP1; 2000-638216/61.
P-PSDB; AAB00184.
                                                                                Gish KC;
                    29-NOV-1999;
                               02-DEC-1999
                                                                                                                                                                                                                                                                                                                                                                          Query Match
                                                                                Mack D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           61
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BCHl genes and protein useful for the diagnosis and treatment of breast
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present sequence encodes a human breast cancer protein, designated
                                                         Nucleotide sequence of a human breast cancer protein designated BCH1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           61 agtoctggtactcttgggagtttccatcttctggtctctgcccagaatccgacaacagc 120
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                                                                                                                     Breast cancer protein; BCH1; breast cancer; chromosome 12; 12q12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ö
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/product= "breast cancer protein BCH1"
47..103
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ouery Match 99.7%; Score 469.4; DB 22; Best Local Similarity 99.8%; Pred. No. 3.4e-119; Matches 470; Conservative 0; Mismatches 1;
                                                                                                                                                                                                                                                                           Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; Fig 1; 73pp; English.
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                                                                                                                                                     anti-oestrogen therapy; ss.
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/*tag=
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                                                                                                                                                                                                               Homo sapiens
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23-JUL-2001
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This sequence encodes a human breast tumour protein immunogenic fragment of the invention. The polypeptides or nucleic acids encoding them are useful in vaccines and pharmaceutical compositions for manufacture of medicaments for inhibiting the development of breast cancer in a patient. They can also be used to treat breast cancer. Antibodies against these polypeptides can be used to detect and monitor progression of breast cancer in patients. Primers and probes derived from the polynucleotides encoding the breast proteins are useful for detection of breast cancer. Peripheral blood cells from a patient incubated in the presence of at least one polypeptide, such that T cells proliferate, are useful in
240
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                                                     auccactyctyctycaacaactycyaccactyctyctcctaccactycaaccaccyctyc
                                                                                                                                                                                                                                                                                                                                                                           protein; immunogenic fragment; vaccine; detection;
development; therapy; ss.
                                                                                                                                                                                                                                                                                                                                                      human breast tumour protein immunogenic fragment.
                                                                                                                                                                                                               New breast tumour protein genes used, or tor diagnosis of breast cancer
                                                                                                                                                                                                                                                                                        ВР
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                                                                                                                                                                                                                                                                                      AAX84202 standard; cDNA; 480
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97US-0998253.
97US-0998255.
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24-DEC-1997
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                                                                                                                                                                                                                                                                    manufacture of a medicament for treating breast cancer in a patient.
Antigen presenting cells incubated in the presence of at least one
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 183 aaccactgctgctgcaaccactgcgaccactgctgctcctaccactgcaaccaccgctgc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             aaccactgctgctgctacaactgcgaccactgctgctcctaccactgcaaccaccgctgc
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                                                                                                                                                                        95.2%; Score 448.4; DB 20; Length 480; 98.5%; Pred. No. 2e-113;
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                                                   polypeptide are also useful for treating breast cancer.
                                                                                              Sequence 480 BP; 121 A; 132 C; 87 G; 140 T; 0 other;
                                                                                                                                                                        Score 448.4,
Pred. No. 2e-113;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CDNA sequence of human breast tumour clone 1016F8
                                                                                                                                                                                                                           0; Mismatches
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                                                                                                                                                                                                                           Conservative
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                                                                                                                                                                                               Similarity
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02-JUL-1999;
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Matches
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endocrine; metabolism; regulation; malabsorption; gastritis; neoplasm

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                                                                                                  The present sequence was isolated from a breast tumour cDNA library. It is provided in a specification relating to compounds for immunotherapy and diagnosis of breast cancer. Breast tumour antigens and the polynucleotides that encode them may be used in the production of a pharmareutical composition to be used in the treatment of breast cancer. Proliferated T cells and incubated antigen presenting cells are also required. The polypeptides and polynucleotides may also be used to
ted polypeptide comprising an immunogenic portion of a protein useful in the detection and treatment of breast
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                                                                 Claim 4; Page 74; 95pp; English.
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                                                                                                                                                                                                                                                                                                                                                Conservative
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Matches 463; Conserv
                                                                                                                                                                                                                           produce a vaccine.
novel isolated
                breast cancer
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                                                                     Human secreted protein gene 34 clone HMQAJ64.
   (first entry)
25-MAR-1999
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New isolated human genes and the secreted polypeptides they encode useful for diagnosis and treatment of e.g. cancers, neurological
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Ni J, Olsen HS;
                                                                                                                                                                                                                                                                                                                                                   useful for diagnosis and treatment of e.g. cancers, neurolo disorders, immune diseases, inflammation or blood disorders
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DW, Moore PA,
, Young P;
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97US-0048094.
97US-0048095.
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97US-0041276.
97US-0041277.
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97US-0048160.
97US-0048186.
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97US-0042344
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                Homo sapiens.
                              WO9842738-A1
                                                              19-MAR-1998;
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30-MAY-1997;
05-AUG-1997;
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                                                                                    Human secreted protein gene 34 clone HMQAJ64.
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secreted human protein. The gene number, and the clone it is derived from, are detailed in the descriptor line. The gene can be used to generate fusion proteins by linking to the gene to a human immunoglobulin by cortion (e.g. AAX00051) for increasing the stability of the fused protein as compared to the human protein only.

The invention relates to 87 novel genes and their fragments (nucleic acid sequences: AAX0611-X0074: amino acid sequences AAX06004) which are useful for preventing, treating or ameliorating medical conditions e.g. by protein or gene therapy. Also, pathological conditions can be diagnosed by determining the amount of the new polypeptides in a sample or by determining the presence of mutations in the new polynucleotides. Specific uses are described for each of the 87 polynucleotides, based on which tissues they are most highly expressed in (see AAX00611 for described uses).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   sequence represents a nucleic acid molecule which encodes a sted human protein. The gene number, and the clone it is derived
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                                                                                                                                                                                                                                                                                                                  Olsen HS;
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                                                                                                                                                                                                                                                                                                               Ni J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 20;
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Pred. No. 2e-113;
); Mismatches (
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Lafleur DW, Moore PA,
Shi Y, Young P;
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97US-0048187.
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Ruben SM,
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This sequence encodes a human secreted protein of the invention. The extended CDNAs (or genomic DNAs obtainable from them) may be used to prepare PCR primers and probes. These are useful for forensic matching or positive identification by DNA sequencing. They may also be used in a lternative fingerprint identification techniques. Antibodies against the proteins encoded by the extended cDNAs are useful in identification of tissue types or cell species, as well as identifying tissue specific soluble proteins. The sequences can be used for chromosome mapping and identification of genes associated with hereditary diseases or drug response. Signal sequences from the cDNAs can be used to clone upstream genomic DNA sequences including promoters. This is in turn useful for identifying proteins that interact with promoter sequences. Some of the proteins may be useful in diagnosing and treating sequences. Some of the proteins may be useful in diagnosing and treating several disorders including, but not limited to: cancer, hyperlipidaemia, cardiovascular and neurodegenerative disorders, autoimmune diseases, and rheumatic diseases, embryogenic disorders, hypertension, renal injury, and monacidurias, hypoglycaemia, male rat infertility and myopathies.
                                                                                                                                                                                                                                                                                      chromosome mapping; human; hereditary disease; diagnosis; cancer; hyperlipidaemia; cardiovascular; neurodegenerative disorder; therapy; autoimmune disease; rheumatic disease; embryogenic disorder; myopathy; renal injury; amino aciduria; hypoglycaemia; male rat infertility; hypertension; ss.
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Secreted protein EST coding sequence 108-002-5-0-F3-FL.
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                                                                                                                             BP.
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98US-0081563.
98US-0096116.
98US-0099273.
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The present sequence was isolated from a breast tumour cDNA library. It is provided in a specification relating to compounds for immunotherapy
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breast
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A novel isolated polypeptide comprising an immunogenic portion breast cancer protein useful in the detection and treatment of
                                                                                                                                                                                                                                                                                   antigen; cytostatic; immunotherapy;
                                                                                                                                                                                                                                                                                                                                                                                                     cDNA sequence of human breast tumour clone B511S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 4; Page 88; 95pp; English.
                                                                                                                                                                                                                                                                                                                                           AAC79470 standard; cDNA; 578
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                                                                                                                                                                                                                                                                                                                                                                                                                        breast tumour antigonancer; vaccine; ss
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Indels

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95.2%; Score 448.4; DB 20; Length 497; 98.5%; Pred. No. 2e-113;

Pred. No. 2e-113; 0; Mismatches

Best Local Similarity 98.5 Matches 463; Conservative

Query Match

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20-JUL-2000; 27-JUL-2000; 218-DEC-2000; 2
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and diagnosis of breast cancer. Breast tumour antigens and the polynucleotides that encode them may be used in the production of a pharmaceutical composition to be used in the treatment of breast cancer. Proliferated T cells and incubated antigen presenting cells are also required. The polypeptides and polynucleotides may also be used to produce a vaccine.
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                                                          C; 88 G; 147 T; 0 other;
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                                                                                 Score 448.4; DB 21;
Pred. No. 2.2e-113;
0; Mismatches 6;
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Similarity 98.5%;
3; Conservative
                                                          Sequence 578 BP; 206 A; 137
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          B511S cDNA sequence
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Best Local Simi
Matches 463;
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The invention relates to identifying tissue-specific polynucleotides (P) from tissue of interest (TI), performing DNA microarray analysis to identify first subset of polynucleotides (SPI) at least 2-fold over expressed in TI, and performing quantitative polymerase chain reaction (PCR) analysis on SPI to identify second subset of (P). The method is useful for determining the presence or absence of a cancer cell in a patient, monitoring the presence or absence of a cancer cell in a piological sample such as blood, serum, lymph nodes, bone marrow, sputum, urine or a tumour blopsy sample. The methods are useful for determining the presence of or monitoring progression of prostate, breast, out, und, head and neck, lymphoma, leukemia, melanoma, liver, squstic, kidhey, bladder, pancreatic or endometrial cancer. The present sequence represents B511S CDNA.
                                                                                                                                                                                                                                                                                            Identifying tissue (tumour)-specific polynucleotides overexpressed in tissue of interest as compared to control tissue, for detecting cancer cells in patient, comprises DNA microarray analysis or quantitative
                                                                                                                                                              Persing DH;
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Pred. No. 2.2
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                                                                                                                                                              Molesh DA,
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2000US-219862P.
2000US-221300P.
2000US-256592P.
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Best Local Similarity 98.5%;
Matches 463; Conservative
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                                                                                                                                                              Dillon
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P-PSDB; AAG65988.
                                                                                                             (CORI-) CORIXA CORP
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Sequences AAX55578-82 represent overlapping polynucleotide sequences of BS106 clones isolated from the breast tissue. The BS106 nucleic acids correspond to a breast tissue gene and can be used as markers for breast tissue disease. The products can be used for detecting, diagnosing, staging, monitoring, prognosticating, preventing or treating, or determining diseases or conditions associated with BS106, especially
318 tggtagagtgtgtccctgagatggaatcagcttgagtcttctgcaatt-ggtcacaacta 376
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                          BS106; breast; marker; detection; diagnosis; breast cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Kratochvil
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                                                                                    Sequence ^82 BP; 109 A; 136 C; 90 G; 145 T; 2 other;
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Pred. No. 4.9e-113;
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Stroupe SD;
                                                                                                                                                                                                                                                                                   BS106 consensus polynucleotide sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New isolated BS106 nucleic acids
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Roberts-rapp L, Russell JC,
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98.9%;
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                                                                                                                                                                                          AAX55581 standard; DNA;
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P-PSDB; AAY13466.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BS106 is a breast tissue gene with which breast cancer and related diseases are associated. The BS106 polynucleotides AAV31989-V31993 can be detected by BS106 specific polynucleotides or complements acting as useful markers for detection methods. The products and methods can be used for detecting, diagnosing, staging, preventing or treating, or determining predisposition to diseases or conditions of the breast such as breast cancer.
                                                                                                                                          human; BS106; breast tissue gene; breast cancer; detection marker.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1;
                                                                                                                                                                                                                                                                                                                                                                                                                                         Detecting BS106 gene products - useful for developing products detecting, staging, preventing, treating or determining predisposition to e.g. breast cancer
                                                                                                                                                                                                                                                                                                                                                        Friedman PN;
MR, Kratochvil JD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               94.9%; Score 447; DB 19; Length 482;
llarity 98.9%; Pred. No. 4.9e-113;
Conservative 2; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 482 BP; 109 A; 136 C; 90 G; 145 T; 2 other;
                                                                                                                                                                                                                                                                                                                                                      M, Colpitts TL,
Hodges SC, Klass
JC, Stroupe SD;
                                                                                                              BS106 polynucleotide consensus sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 1; Page 88; 114pp; English
                          DNA; 482 BP.
                                                                                                                                                                                                                                                                                                                                                                    Granados EN, Hodo
pp L, Russell JC,
                                                                                                                                                                                                                                                               97WO-US19836
                                                                                                                                                                                                                                                                                                                                                      Cohen M,
                                                                                                                                                                                                                                                                                            96US-0742067
                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 1998-272234/24.
                        AAV31992 standard;
                                                                                                                                                                                                                                                                                                                         (ABBO ) ABBOTT LAB.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
                                                                                                                                                                                                                                                                                                                                                      Billing-Medel PA,
                                                                                                                                                                                                                                                                                                                                                                    Gordon J, Granad
Roberts-Rapp L,
                                                                                                                                                                         Homo sapiens
                                                                                                                                                                                                     WU9818945-A1
                                                                                                                                                                                                                                                             31-0C1-1997;
                                                                                                                                                                                                                                                                                            31-OCT-1996;
                                                                                  25-SEP-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Simi
Matches 458;
                                                                                                                                                                                                                                 07-MAY-1998.
                                                    AAV31992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     6.1
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Gordon
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                                                                                                                                                                                                                                                                                                            ss; human; BS106; breast tissue gene; breast cancer; detection marker.
tggtagagtgtgtccctgagatggaatcagcttgagtcttctgcaattgggtcacaacta 360
                                                                                        for
                                       Ltctaccactgctcgtaaagacattccagttttacccaaatgggttggggatctcccgaa
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               Kratochvil JD;
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MR, Kratoch
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 553 BP; 159 A; 144 C; 101 G; 148 T; 1 other;
                                                                                                                                                          Colpitts TL,
les SC, Klass
                                                                                                                                                                                                                                                                                         BS106 polynucleotide clone 1662885 (ii).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Stroupe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 1; Page 90; 114pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Cohen M, Col
s EN, Hodges
                                                                                                                                                                                                                              ВР
                                                                                                                                                                                                                             AAV31993 standard; DNA; 553
                                                                                                                                                                                                                                                                                                                                                                                          97WO-US19836
                                                                                                                                                                                                                                                                                                                                                                                                              96US-0742067
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Russell
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Granados EN,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1998-272234/24.
                                                                                                                                                                                                                                                                                                                                                                                                                                  (ABBO ) ABBOTT LAB.
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Billing-Medel PA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     breast cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Roberts-Rapp L,
                                                                                                                                                                                                                                                                                                                                                   WO9818945-A1
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DB 19; Length 553; Indels

Score 446.8; DB 19; Pred. No. 5.9e-113; 0; Mismatches 7;

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Conservative

462;

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Similarity

Query Match Best Local Si Matches 462;

94.9%; 98.3%;

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Sequences AAX5558-82 represent overlapping polynucleotide sequences of BS106 clones isolated from the breast tissue. The BS106 nucleic acids correspond to a breast tissue gene and can be used as markers for breast tissue disease. The products can be used for detecting, diagnosing, staging, monitoring, prognosticating, preventing or treating, or determining diseases or conditions associated with BS106, especially
                                                                                    180
                                                                                                                                                                                                                                      360
                                                            151
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                                                                                                                                                                                                                                                                                                              450
                                   tgctccagctgacacgtatccagctactggtcctgctgatgatgaagcccctgatgatgctga
                                                                                                                                                                                                              ttctaccactgctcgtaaagacattccagttttacccaaatgggttggggatctcccgaa
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                                                                                                                                                                                                                                                   tgctccagctgacacgtatccagctactggtcctgctgatgatgaagcccctgatgctga
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                                                                                                                                                                                                                                                                                                                                      Friedman PN;
Kratochvil
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Colpitts TL,
SC, Klass M
Stroupe SD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BS106 consensus polynucleotide sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                             BP.
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Russell JC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Cohen M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               98WO-US22020
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                                                                                                                                                                                                                                                                                                                                                                                                                            AAX55582 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Granadose,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1999-326991/27
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New isolated BS106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Billiing-Medel PA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Roberts-rapp L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      P-PSDB; AAY13466
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO9923230-A1
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This sequence represents a breast specific gene (BSG) clone. The BSG sequences are also referred to as breast specific markers (BSM). The invention relates to methods for diagnosing, monitoring, staging, imaging and treating breast cancer (BC). The methods comprise measuring the levels of BSG products in cells, tissues or body fluids of the patient and comparing the measured levels of BSG, with BSG levels of a normal human control. An antibody against the BSG sequences can be labelled and used for imaging BC in a patient. The antibody can be conjugated to a cytotoxic agent, and used for treating BC in a patient.
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                                                                                                                                                                                                                                                                                                                                                                                                                 Human BS106 gene-specific full length cDNA, EST clone 1662885inh.
                                                                                                                                                                                     ;
                                                                                                                                                                  Length 544;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 9S106 protein; breast cancer; metastasis; gene therapy; expressed sequence tag; EST; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                     Sequence 544 BP; 142 A; 143 C; 94 G; 154 T; 11 other;
                                                                                                                                                                 DB 21;
                                                                                                                                                                                     9
                                                                                                                                                                 Score 446.4; DB 21
Pred. No. 7.5e-113;
0; Mismatches 6;
                    7; Page 38; 45pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ВР
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     standard; cDNA; 472
                                                                                                                                                                 Ouery Match 94.8%;
Best Local Similarity 98.5%;
Matches 461; Conservative
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                    Claim
   genes
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AAD18673
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                                                                                                                                                                                                                                           tygtagagtgtgtccctgagatggaatcagcttgagtcttctgcaattgggtcacaacta 360
                                                                                                                                                                                                                                                                                            212 aaccactyctyctycaaccactycyaccactyctyctcctaccactycaaccaccyctyc 271
                                                                                                                                                                                                                                                                                                                       ttcatgcttcctgtgatttratccaactacttaccttgcctacgatatcccctttatctc 420
                                                                   Gaps
                                                                                   Detecting, diagnosing, monitoring, staging, prognosticating, imaging and treating breast cancer using protein product of breast specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           specific gene; breast specific marker; BSG; diagnosis;
                                                                 1;
                                                                                                                                                                                                                                                                                                                                                            20; Length
                                                                  Indels
                  G; 148 T; 1 other;
                                              Score 446.8; DB 20;
Pred. No. 5.9e-113;
3; Mismatches 7;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human breast specific gene clone Mam002
                  Sequence 553 BP; 159 A; 144 C; 101
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      therapy; Mam002; ss
                                                                 0;
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                                              94.9%;
98.3%;
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                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                               AA291765 standard; DNA;
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                                                         Similarity
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 breast cancer.
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                                                                 462;
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                                               Ouery Match
                                                         Best Local
Matches 46
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                                                                                                                                                gene or its fragment which codes for a BS106 polypeptide, useful for he detection of a breast disease such as breast cancer -
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Billing-Medel PA, Cohen M, Colpitts TL, Friedman PN, Gordon J;
Granados EN, Hodges SC, Klass MR, Kratochvil JD, Roberts-Rapp L;
Russell JC, Stroupe SD;
                                                                                                                                                                                                                                                                                                                                                                                                                  1; Gaps
                                                                                                                                                                                                         The Invention relates to human BS106-specific polypeptides and polymucleotides. The BS106 polypeptides and antibodies are useful for detecting, diagnosing, staging, monitoring, prognosticating, preventing, treating or determining the predisposition of an individual to diseases and conditions of the breast such as breast canner. They are also useful in the treatment of tumours or metastases. Polynucleotides of the invention are useful in drug screening and gene therapy. The present sequence is human BS106 qene specific full length CDNA, expressed sequence tag (EST) clone 1662885inh.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       tqctccagctgacacgtatccagctactgqtcctgctgatgatgaagcccctgatgctga
                                                                                                                                                                                                                                                                                                                                                                                    Query Match

94.7%; Score 446.2; DB 22; Length 472;
Best Local Similarity 99.1%; Pred. No. 8e-113;
Matches 459; Conservative 0; Mismatches 3; Indels 1;
                                                                                                                                                                                                                                                                                                                                                 Sequence 472 BP; 108 A; 134 C; 86 G; 144 T; 0 other;
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                                                                                                                                                                                   Example 1; Fig 1; 158pp; English.
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Search completed: October 5, 2002, 05:44:20 Job time: 3916 sec

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Sequence 41
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                                                                           Complementary DNAs
Patent: US 6312922-A 41 06-NOV-2001;
Location/Qualifiers
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Edwards, J.-B. Dumas. Milne., Duclert, A.
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                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Metazoa;
Mammalia; Eutheria;
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                                                                                                                                Similarity
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1 (bases 1 to 482)

Medel, P.A.B., Cohen, M., Colpitts, T.L., .

Granados, E.N., Hodges, S.C., Klass, M.R., .

Russell, J.C. and Stroupe, S.D.

Patent: JP 2001503980-A 4 27-MAR-2001;

ABBOTT LABORATORIES

OS Unidentified

PN JP 2001503980-A/4

PD 27-MAR-2001

PF 31-OCT 1997 -

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tch 94.9%; al Similarity 98.9%; 458; Conservative
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STEVEN D STROUPE
PC C12N15/09, 6
PC C07K16/30,
PC C12N5/10,C12
A61K37/02
CC Strandedness
CC Topology: Li
FH Key
FT source
FT
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PI
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31-OCT-1996 US 08/742067
PATRICIA A BILLING MEDEL, MAI
N FRIEDMAN,
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                                                                                                                                                                                              Strandedness:
                                                                                                                                                                                                                                                                      JOHN D KRATOCHVIL, LISA ROBERTS
                                                                             Ω
                                                                            /organism="unidentified"
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Score 447; DB Pred. No. 3.5e 2; Mismatches
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                                                                                                                                                                                                                                                                                                                       MEDEL, MAURICE COHEN, TRACY L COLPITTS.
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OS Unidentified
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PN JP 2001503980-A/5
PD 27-MAR-2001
PF 31-OCT-1996 US 08/742067
PR 31-OCT-1996 US 08/742067
PR 31-OCT-1996 US 08/742067
PI PATRICIA A BILLING MEDEL, MAURICE COHEN, TRACY L COLPITTS, PI
POLA N FRIEDMAN,
PI JULIAN GORDON, EDWARD N GRANADOS, STEVEN C HODGES, PI MICHAEL
R KLASS,
PI JOHN D KRATOCHVIL, LISA ROBERTS RAPP, JOHN C RUSSELL, PI
STEVEN D STROUPE
C C12N15/09, A61K38/00, A61K39/395, A61K39/395, A61P35/00, C07K14/47,
PC C12N15/09, A61K38/00, A61K39/395, A61R39/395, A61P35/00, C07K14/47,
PC C12N5/10, C12P21/02, C12Q1/68, G01N33/53, C12N15/00, C12N5/00, PC
A61K37/02
CC Strandedness: Single:
CC Strandedness: Single:
FH Key
1. 553
FT source
//organism='Unidentified'.
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Reagent and method useful in detecting mammary diseases Patent: JP 2001503980-A 5 27-MAR-2001;
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Sequence 5 from Patent WO0165262.
AX240679
AX240679.1 GI:15797662
                                                                                                                                                                                    Eukaryota; Metazoa; Chordata; Craniata; Vertebrala; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; I (bases | to 472)
Billing-Medel, P.A., Cohen, M., Colpitts, T.L., Friedma Gordon, J., Granados, E.N., Hodges, S.C., Klass, M.R., K Roberts-Rapp, L., Russell, J.C. and Stroupe, S.D. Raagents and methods useful for detecting diseases c Patent; WO 0165262-A 5 07-SEP-2001;
                              Similarity
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/db_xref="taxon:32644"
a 144 c 101 g 141
                                                                                                                                                           Location/Qualifiers
                              94.78;
99.18;
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             Score 446.2; DB b;
Pred. No. 5.4e-96;
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ACCESSION VERSION KEYWORDS SOURCE

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AUTHORS

COMMENT

Friedman, P.N

Euteleustomi;

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Kratochvil, J.D.

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tgctccagctgacacgtatccagctactggtcctgctgatgatgatgaagcccctgatgctga
                               TAATCAGTTTATTTTCTTTCAAATAAAAAAATAACTATGAGCAA 470
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ABBOTT LABORATORIES (US)
Location/Qualifiers
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AX240680
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Location/Qualifiers
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1 from Patent WO0135811.
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/db_xref="taxon:9606"
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455; Conserv
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Sequence 26
Ax067322
Ax067322.1
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/db_xref="taxon:9606"
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Primates;
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Pred. No. 4.9e-95;
2; Mismatches 5
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Best Local Similarity
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Miksicek,R.J., Myal,Y.,
Direct Submission
Submitted (23-AUG-2001)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens small breast AF414087 AF414087.1 GI:15559110
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       gene
                                                                                                                                                                                                                                                                                                                       /product="small breast epithelial mucin"
/protein_id="AALI02119_1"
/db_xref="GI:1559111"
/translation="MRELAVLYLLGVSIFLVSAQNPTTAAPANTYPATGPANDEAPDA
ETTAAATTATTAAPTTATTAASTTARKDIPVLPKWVGDLPNGKVCP"
a 117 c 80 g 116 t
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47. .319
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/codon_start=1
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/db_xref="taxon:9606"
/chromosome="12"
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/note="corresponds to SAGE tag: cttcctgtga
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Pred. No. 4.8e-81
0; Mismatches
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Bannatyne Av
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Sequence 2 from F
AX240676
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                                                  BD007429.1 GI:18635800
JP 2001503980-A/2.
unidentified.
  1 (bases 1 to 308)
Medel, P.A.B., Cohen
                                      unidentified
                                                                                     Reagent and method useful BD007429
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                           unclassified
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99.3%;
  Cohen, M., Colpitts, T.L.,
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Patent WO0165262
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  Friedman, P.N.,
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  Gordon, J.,
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synthetic construct.
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artificial sequence.
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                                                                      Sequence 3
AX240677
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C C12N15/09,A61K38/00,A61K39/395,A61K39/395,A61P35/00,C07K14/47,
PC C07K16/30,
PC C12N5/10,C12P21/02,C12Q1/68,G01N33/53,C12N15/00,C12N5/00, PC
A61K37/02

CC Strandedness: Single;
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R KLASS,
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POLA N FRIEDMAN,
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AX328499
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammaila; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                               Mack,D. and Gish,K.C.

Novel methods of diagnosing and determining prognosis of breast cancer, compositions, and methods of screening for breast cance
                                                                                                                                                                      Patent: WO 0135811-A 2 25-MAY-2001; EOS Biotechnology, Inc. (US)
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Baker,K.P., Goddard,A. and Wood,W.I.
Human polypeptides and methods for the use thereof Patent: WO 0107611-A 220 01-FEB-2001;
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 266)
Baker,K.P., Goddard,A. and Wood,W.I.
Human polypeptides and methods for the use thereof
Patent: WO 0107611-A 219 01-FEB-2001;
Genentech, Inc. (US)
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